
Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=6; day=18; hr=16; min=42; sec=8; ms=656;]

Validated By CRFValidator v 1.0.3

Application No: 10574424 Version No: 3.1

Input Set:

Output Set:

Started: 2008-06-18 16:37:56.814

Finished: 2008-06-18 16:37:58.167

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 353 ms

Total Warnings: 12

Total Errors: 0

No. of SeqIDs Defined: 12

Actual SeqID Count: 12

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W	402	Undefined organism found in <213> in SEQ ID (3)
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W	402	Undefined organism found in <213> in SEQ ID (5)
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W	402	Undefined organism found in <213> in SEQ ID (8)
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Ala	Ala	Thr 35	Val	Ala	Ile	Ala	Ala 40	Ala	Tyr	Asn	Asn	Gly 45	Gln	Glu	Ile
Asn	Gly 50	Phe	Lys	Ala	Gly	Glu 55	Thr	Ile	Tyr	Asp	Ile 60	Asp	Glu	Asp	Gly
Thr 65	Ile	Thr	Lys	Lys	Asp 70	Ala	Thr	Ala	Ala	Asp 75	Val	Glu	Ala	Asp	Asp
Phe	Lys	Gly	Leu	Gly 85	Leu	Lys	Lys	Val	Val 90	Thr	Asn	Leu	Thr	Lys 95	Thr
Val	Asn	Glu	Asn 100	Lys	Gln	Asn	Val	Asp 105	Ala	Lys	Val	Lys	Ala 110	Ala	Glu
Ser	Glu	Ile 115	Glu	Lys	Leu	Thr	Thr 120	Lys	Leu	Ala	Asp	Thr 125	Asp	Ala	Ala
Leu	Ala 130	Asp	Thr	Asp	Ala	Ala 135	Leu	Asp	Ala	Thr	Thr 140	Asn	Ala	Leu	Asn
Lys 145	Leu	Gly	Glu	Asn	Ile 150	Thr	Thr	Phe	Ala	Glu 155	Glu	Thr	Lys	Thr	Asn 160

Ile Val	Lys	Ile	Asp 165	Glu	Lys	Leu	Glu	Ala 170	Val	Ala	Asp	Thr	Val 175	Asp
Lys His	Ala	Glu 180	Ala	Phe	Asn	Asp	Ile 185	Ala	Asp	Ser	Leu	Asp 190	Glu	Thr
Asn Thr	Lys 195	Ala	Asp	Glu	Ala	Val 200	Lys	Thr	Ala	Asn	Glu 205	Ala	Lys	Gln
Thr Ala	Glu	Glu	Thr	Lys	Gln 215	Asn	Val	Asp	Ala	Lys 220	Val	Lys	Ala	Ala
Glu Thr 225	Ala	Ala	Gly	Lys 230	Ala	Glu	Ala	Ala	Ala 235	Gly	Thr	Ala	Asn	Thr 240
Ala Ala	Asp	Lys	Ala 245	Glu	Ala	Val	Ala	Ala 250	Lys	Val	Thr	Asp	Ile 255	Lys
Ala Asp	Ile	Ala 260	Thr	Asn	Lys	Asp	Asn 265	Ile	Ala	Lys	Lys	Ala 270	Asn	Ser
Ala Asp	Val 275	Tyr	Thr	Arg	Glu	Glu 280	Ser	Asp	Ser	Lys	Phe 285	Val	Arg	Ile
Asp Gly 290	Leu	Asn	Ala	Thr	Thr 295	Glu	Lys	Leu	Asp	Thr 300	Arg	Leu	Ala	Ser
Ala Glu 305	Lys	Ser	Ile	Ala 310	Asp	His	Asp	Thr	Arg 315	Leu	Asn	Gly	Leu	Asp 320
Lys Thr	Val	Ser	Asp 325	Leu	Arg	Lys	Glu	Thr 330	Arg	Gln	Gly	Leu	Ala 335	Glu
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Thr Ile	Tyr 35	Asp	Ile	Asp	Glu	Asp 40	Gly	Thr	Ile	Thr	Lys 45	Lys	Asp	Ala
Thr Ala	Ala	Asp	Val	Glu	Ala 55	Asp	Asp	Phe	Lys	Gly 60	Leu	Gly	Leu	Lys
Lys Val 65	Val	Thr	Asn	Leu 70	Thr	Lys	Thr	Val	Asn 75	Glu	Asn	Lys	Gln	Asn 80

Val Asp A	la Lys	Val 85	Lys	Ala	Ala	Glu	Ser 90	Glu	Ile	Glu	Lys	Leu 95	Thr
Thr Lys L	eu Ala 100	Asp	Thr	Asp	Ala	Ala 105	Leu	Ala	Asp	Thr	Asp 110	Ala	Ala
Leu Asp A 1	la Thr 15	Thr	Asn	Ala	Leu 120	Asn	Lys	Leu	Gly	Glu 125	Asn	Ile	Thr
Thr Phe A	la Glu	Glu	Thr	Lys 135	Thr	Asn	Ile	Val	Lys 140	Ile	Asp	Glu	Lys
Leu Glu A 145	la Val	Ala	Asp 150	Thr	Val	Asp	Lys	His 155	Ala	Glu	Ala	Phe	Asn 160
Asp Ile A	la Asp	Ser 165	Leu	Asp	Glu	Thr	Asn 170	Thr	Lys	Ala	Asp	Glu 175	Ala
Val Lys T	hr Ala 180	Asn	Glu	Ala	Lys	Gln 185	Thr	Ala	Glu	Glu	Thr 190	Lys	Gln
Asn Val A 1	sp Ala 95	Lys	Val	Lys	Ala 200	Ala	Glu	Thr	Ala	Ala 205	Gly	Lys	Ala
Glu Ala A 210	la Ala	Gly	Thr	Ala 215	Asn	Thr	Ala	Ala	Asp 220	Lys	Ala	Glu	Ala
Val Ala A 225	la Lys	Val	Thr 230	Asp	Ile	Lys	Ala	Asp 235	Ile	Ala	Thr	Asn	Lys 240
Asp Asn I	le Ala	Lys 245	Lys	Ala	Asn	Ser	Ala 250	Asp	Val	Tyr	Thr	Arg 255	Glu
Glu Ser A	sp Ser 260	Lys	Phe	Val	_	Ile 265	_	Gly	Leu	Asn	Ala 270	Thr	Thr
Glu Lys L	eu Asp 75	Thr	Arg	Leu	Ala 280	Ser	Ala	Glu	Lys	Ser 285	Ile	Ala	Asp
His Asp T	hr Arg	Leu	Asn	Gly 295	Leu	Asp	Lys	Thr	Val 300	Ser	Asp	Leu	Arg
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20 25 30

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Lys Gly Tyr Thr Pro Gln Ile Ser Val Val Gly Tyr Asn Arg Hi 50 55 60	s Leu
Leu Leu Leu Gly Gln Val Ala Thr Glu Gly Glu Lys Gln Phe Va	1 Gly 80
Gln Ile Ala Arg Ser Glu Gln Ala Ala Glu Gly Val Tyr Asn Ty 85 90 95	r Ile
Thr Val Ala Ser Leu Pro Arg Thr Ala Gly Asp Ile Ala Gly As 100 105 110	p Thr
Trp Asn Thr Ser Lys Val Arg Ala Thr Leu Leu Gly Ile Ser Pr 115 120 125	o Ala
Thr Gln Ala Arg Val Lys Ile Val Thr Tyr Gly Asn Val Thr Ty 130 135 140	r Val
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Ala Thr Tyr Lys Val Asp Glu Tyr His Ala Asn Ala Arg Phe Ala 1 10 15 15 15 15 15 16 16 16 17 17 18 18 18 18 18 18 18 18 18 18 18 18 18	u Thr p Ile r Asp
Ala Thr Tyr Lys Val Asp Glu Tyr His Ala Asn Ala Arg Phe Ala 1 10 15 15 15 15 15 16 16 16 16 16 16 16 16 16 16 16 16 16	u Thr p Ile r Asp p Ile 80

Lys Ala Glu Lys Phe Asn Cys Tyr Gln Ser Pro Met Glu Lys Thr Glu 120 115 Val Cys Gly Gly Asp Phe Ser Thr Thr Ile Asp Arg Thr Lys Trp Gly 130 135 140 Met Asp Tyr Leu Val Asn Val Gly Met Thr Lys Ser Val Arg Ile Asp 150 155 160 Ile Gln Ile Glu Ala Ala Lys Gln 165 <210> 6 <211> 464 <212> PRT <213> N. meningitidis <400> 6 Ser Pro Asp Val Lys Ser Ala Asp Thr Leu Ser Lys Pro Ala Ala Pro 10 Val Val Ser Glu Lys Glu Thr Glu Ala Lys Glu Asp Ala Pro Gln Ala 20 25 30 Gly Ser Gln Gly Gln Gly Ala Pro Ser Ala Gln Gly Ser Gln Asp Met 35 40 Ala Ala Val Ser Glu Glu Asn Thr Gly Asn Gly Gly Ala Val Thr Ala 55 Asp Asn Pro Lys Asn Glu Asp Glu Val Ala Gln Asn Asp Met Pro Gln 75 70 Asn Ala Ala Gly Thr Asp Ser Ser Thr Pro Asn His Thr Pro Asp Pro 90 Asn Met Leu Ala Gly Asn Met Glu Asn Gln Ala Thr Asp Ala Gly Glu 100 105 Ser Ser Gln Pro Ala Asn Gln Pro Asp Met Ala Asn Ala Ala Asp Gly 115 120 125 Met Gln Gly Asp Asp Pro Ser Ala Gly Gln Asn Ala Gly Asn Thr 130 135 Ala Ala Gln Gly Ala Asn Gln Ala Gly Asn Asn Gln Ala Ala Gly Ser 145 150 155 Ser Asp Pro Ile Pro Ala Ser Asn Pro Ala Pro Ala Asn Gly Gly Ser 165 170 Asn Phe Gly Arg Val Asp Leu Ala Asn Gly Val Leu Ile Asp Gly Pro 185 Ser Gln Asn Ile Thr Leu Thr His Cys Lys Gly Asp Ser Cys Ser Gly

195 200 205

Asn Asn Phe Leu Asp Glu Glu Val Gln Leu Lys Ser Glu Phe Glu Lys 215 220 Leu Ser Asp Ala Asp Lys Ile Ser Asn Tyr Lys Lys Asp Gly Lys Asn 230 235 Asp Lys Phe Val Gly Leu Val Ala Asp Ser Val Gln Met Lys Gly Ile 245 250 Asn Gln Tyr Ile Ile Phe Tyr Lys Pro Lys Pro Thr Ser Phe Ala Arg 260 265 Phe Arg Arg Ser Ala Arg Ser Arg Ser Leu Pro Ala Glu Met Pro 280 Leu Ile Pro Val Asn Gln Ala Asp Thr Leu Ile Val Asp Gly Glu Ala 295 300 Val Ser Leu Thr Gly His Ser Gly Asn Ile Phe Ala Pro Glu Gly Asn 315 Tyr Arg Tyr Leu Thr Tyr Gly Ala Glu Lys Leu Pro Gly Gly Ser Tyr 325 330 335 Ala Leu Arg Val Gln Gly Glu Pro Ala Lys Gly Glu Met Leu Ala Gly 340 345 Ala Ala Val Tyr Asn Gly Glu Val Leu His Phe His Thr Glu Asn Gly 355 360 Arg Pro Tyr Pro Thr Arg Gly Arg Phe Ala Ala Lys Val Asp Phe Gly 375 Ser Lys Ser Val Asp Gly Ile Ile Asp Ser Gly Asp Asp Leu His Met 390 395 Gly Thr Gln Lys Phe Lys Ala Ala Ile Asp Gly Asn Gly Phe Lys Gly 410 405 Thr Trp Thr Glu Asn Gly Ser Gly Asp Val Ser Gly Lys Phe Tyr Gly Pro Ala Gly Glu Val Ala Gly Lys Tyr Ser Tyr Arg Pro Thr Asp 440 435 Ala Glu Lys Gly Gly Phe Gly Val Phe Ala Gly Lys Lys Glu Gln Asp 450 455 460 <210> 7 <211> 644 <212> PRT <213> N. meningitidis <400> 7

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Pro	Gln	Asn	Ala	Ala 85	Asp	Thr	Asp	Ser	Leu 90	Thr	Pro	Asn	His	Thr 95	Pro
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Gly	Glu	Ser 115	Glu	Gln	Pro	Ala	Asn 120	Gln	Pro	Asp	Met	Ala 125	Asn	Thr	Ala
Asp	Gly 130	Met	Gln	Gly	Asp	Asp 135	Pro	Ser	Ala	Gly	Gly 140	Glu	Asn	Ala	Gly
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Gly	Ser	Gln	Asn	Pro 165	Ala	Ser	Ser	Thr	Asn 170	Pro	Ser	Ala	Thr	Asn 175	Ser
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Gly	Pro	Ser 195	Gln	Asn	Ile	Thr	Leu 200	Thr	His	Cys	Lys	Gly 205	Asp	Ser	Cys
Ser	Gly 210	Asn	Asn	Phe	Leu	Asp 215	Glu	Glu	Val	Gln	Leu 220	Lys	Ser	Glu	Phe
Glu 225	Lys	Leu	Ser	Asp	Ala 230	Asp	Lys	Ile	Ser	Asn 235	Tyr	Lys	Lys	Asp	Gly 240
Lys	Asn	Asp	Gly	Lys 245	Asn	Asp	Lys	Phe	Val 250	Gly	Leu	Val	Ala	Asp 255	Ser
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